



A GPU-Accelerated Model of Neuroblastoma to Predict Disease Outcome and Find Drug Targets

Kenneth Y. Wertheim, Robert Chisholm, Paul Richmond, Dawn Walker

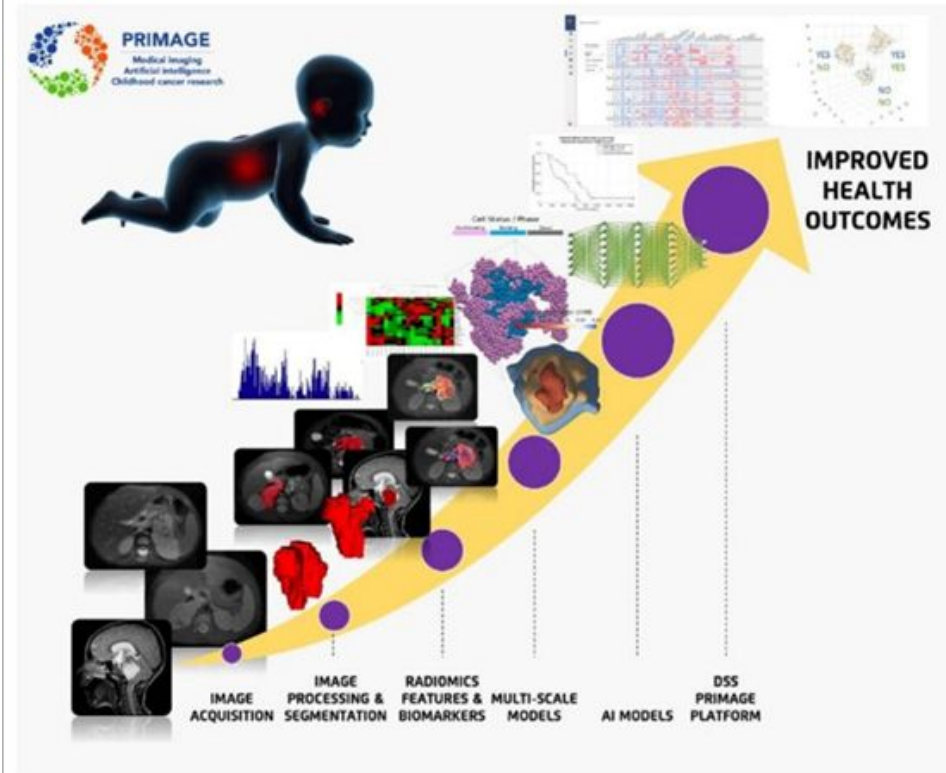
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Horizon 2020
European Union Funding
for Research & Innovation

Objectives

1. PRIMAGE project.
2. Neuroblastoma.
3. First multicellular model of neuroblastoma.
4. Calibration.
5. Clonal competition.
6. Targeted therapies.



Decision support system for the clinical management of malignant solid tumours.

PRIMAGE project: predictive *in silico* multiscale analytics to support childhood cancer personalised evaluation empowered by imaging biomarkers



Luis Martí-Bonmatí^{1*}, Ángel Alberich-Bayarri², Ruth Ladenstein³, Ignacio Blanquer⁴, J. Damian Segrelles⁴, Leonor Cerdá-Alberich⁵, Polyxeni Gkontra⁵, Barbara Hero⁶, J. M. García-Aznar^{7,8}, Daniel Keim⁹, Wolfgang Jentner⁹, Karine Seymour¹⁰, Ana Jiménez-Pastor², Ismael González-Valverde², Blanca Martínez de las Heras¹¹, Samira Essiaf¹², Dawn Walker¹³, Michel Rochette¹⁴, Marian Bubak¹⁵, Jordi Mestres¹⁶, Marco Viceconti¹⁷, Gracia Martí-Besa⁵, Adela Cañete¹¹, Paul Richmond¹³, Kenneth Y. Wertheim¹³, Tomasz Gubala¹⁵, Marek Kasztelnik¹⁵, Jan Meizner¹⁵, Piotr Nowakowski¹⁵, Salvador Gilpérez¹⁸, Amelia Suárez¹⁸, Mario Aznar¹⁸, Giuliana Restante¹⁹ and Emanuele Neri¹⁹

I contributed the first multicellular model of neuroblastoma to the project.

This talk is about what I did with the model outside PRIMAGE.

Martí-Bonmatí, Luis, et al. "PRIMAGE project: predictive in silico multiscale analytics to support childhood cancer personalised evaluation empowered by imaging biomarkers." *European radiology experimental* 4.1 (2020): 1-11.

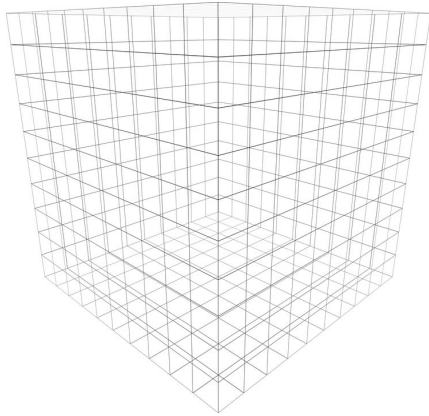
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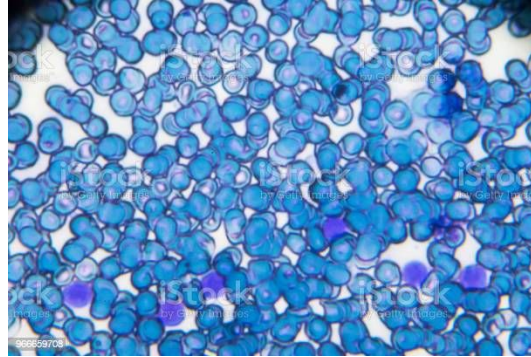
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Multicellular model



Continuous automaton to voxelate the microenvironment.

1. Spatial distributions of cells and extracellular matrix.
2. Concentration dynamics of drugs and nutrients (uniform).

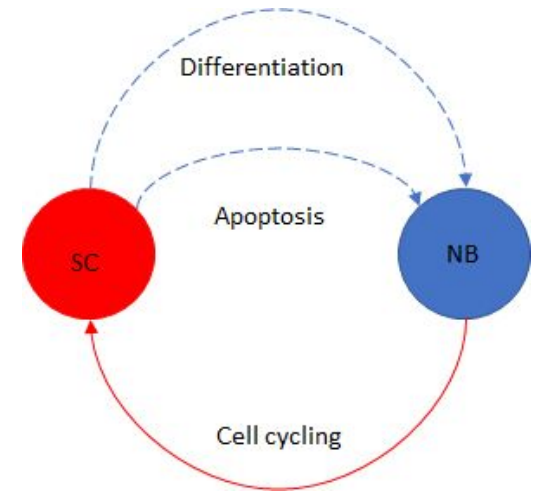


Discrete agents.

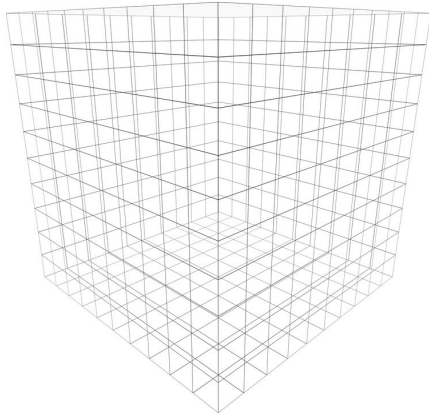
1. Neuroblasts and Schwann cells.
2. Cell cycling and death.

Agent attributes.

1. Mutations.
2. DNA status.
3. Gene expression levels.

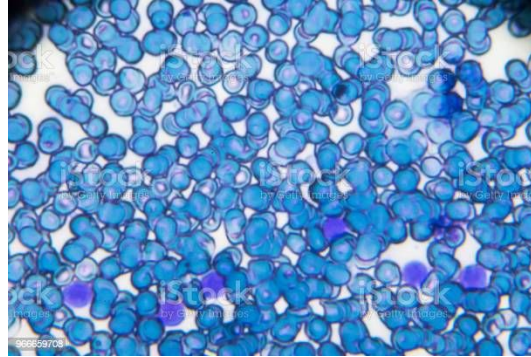


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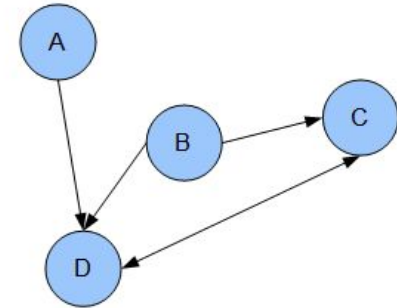
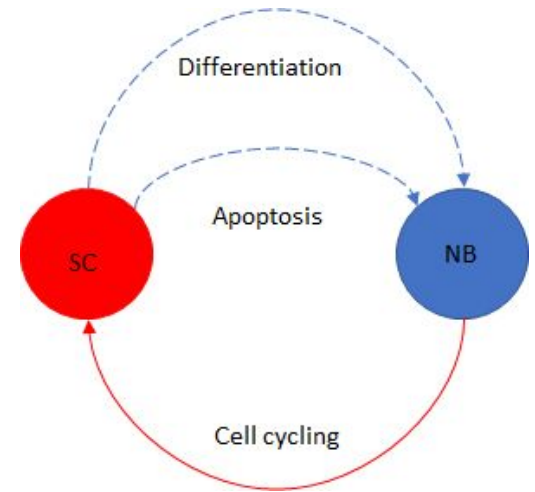


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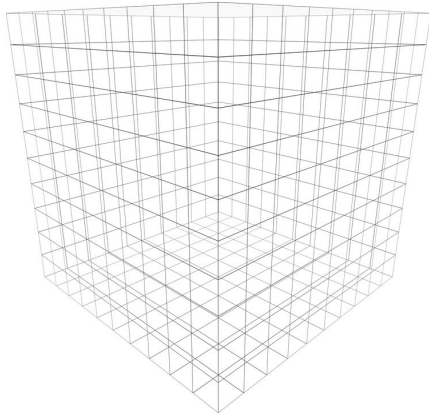
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3. **Gene expression levels.**



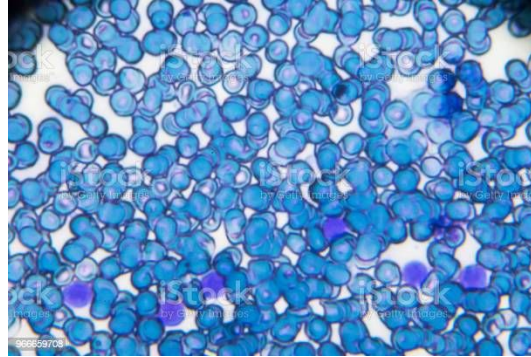
20 gene products. Telomerase, ALT, MYCN, MAPK/RAS pathway, JAB1, CHK1, CDS1, CDC25C, ID2, IAP2, HIF, BNIP3, VEGF, p53, p73, p21, p27, Bcl-2/Bcl-xL, BAK/BAX, and CAS.

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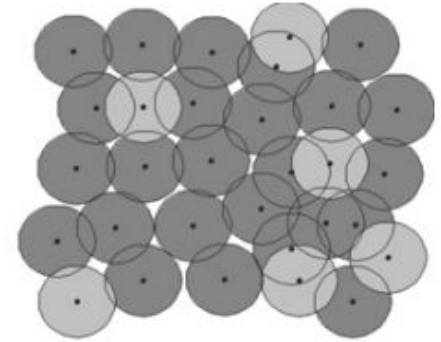


Discrete agents.

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Centre-based mechanical model.

1. Resolve agent-agent overlap and contact inhibition.
2. Linear force law.
3. Equation of motion.

Stochastic simulation algorithm

1. Each agent senses the microenvironment and its neighbouring agents, modifies its behaviour, and updates its attributes.
2. Resolve agent-agent overlap using the mechanical model.
3. Modify the microenvironment by considering the agents collectively.
4. Back to step 1.

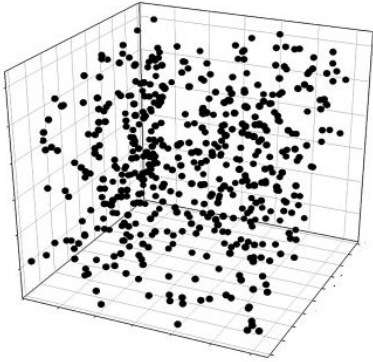
A series of Bernoulli trials. For example, is the MAPK/RAS pathway active?



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Calibration

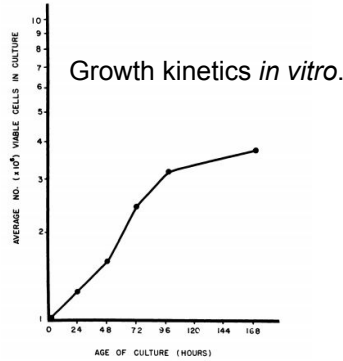


Latin hypercube sampling.

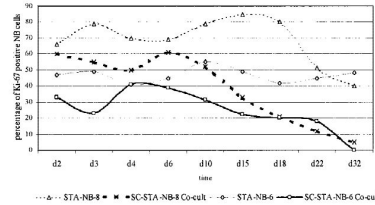
1. 3000 combinations of 20 fitting parameters.

2. Minimised differences between simulation results and *in vitro* data.

3. Refined calibrated parameters for *in vivo* use.



Tumilowicz, Joseph J., et al. "Definition of a continuous human cell line derived from neuroblastoma." *Cancer research* 30.8 (1970): 2110-2118.



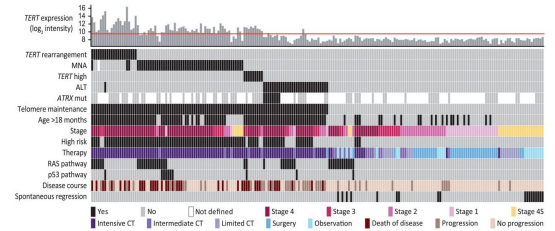
Ambros, Ingeborg M., et al. "Neuroblastoma cells provoke Schwann cell proliferation *in vitro*." *Medical and Pediatric Oncology: The Official Journal of SIOP—International Society of Pediatric Oncology (Société Internationale d'Oncologie Pédiatrique)* 36.1 (2001): 163-168.

Interactions between neuroblastic and Schwann cells *in vitro*.

	Three-stage fit	95% CI	Direct fit	95% CI
Maximum oxygen consumption rate, q_{max} (mmHg · s ⁻¹)	17.5	15.3–25.1	16.3	15.3–17.9
P_{O_2} for 50% drop in consumption, $P_{50,q}$ (mmHg)	2.7	0.0–12.5	1.6	1.2–2.1
Maximum misonidazole binding rate, $k_{b,0}$ ($\times 10^{-4}$ s ⁻¹)	4.5	3.9–4.9	4.4	2.5–5.3
P_{O_2} for 50% drop in binding, $P_{50,b}$ (mmHg)	1.4	0.3–2.6	1.4	1.1–2.5
P_{O_2} for 50% necrosis, $P_{50,n}$ (mmHg)	1.2	0.1–4.9	1.0	0.4–1.2

Warren, Daniel R., and Mike Partridge. "The role of necrosis, acute hypoxia and chronic hypoxia in 18F-FMISO PET image contrast: a computational modelling study." *Physics in Medicine & Biology* 61.24 (2016): 8596.

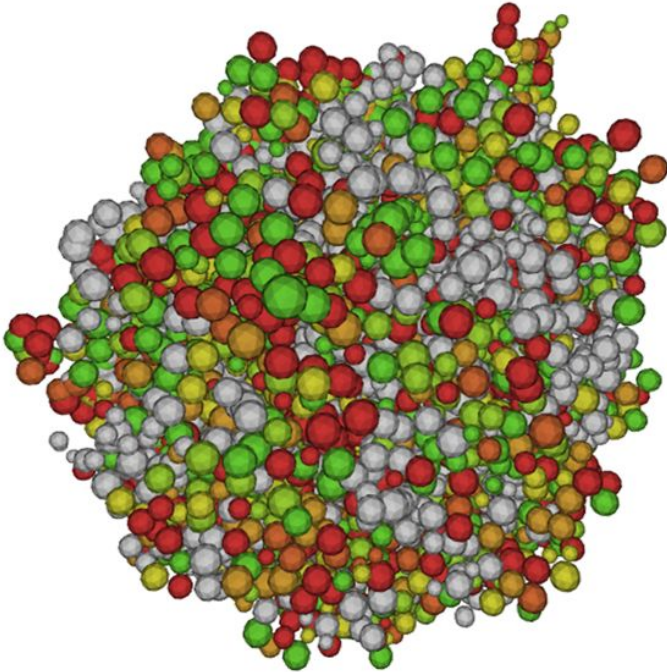
Extent of necrosis during hypoxia *in vitro*.



Ackermann, Sandra, et al. "A mechanistic classification of clinical phenotypes in neuroblastoma." *Science* 362.6419 (2018): 1165-1170.

Clinical outcomes associated with different mutations.

Calibration



Costly simulations.

1. Millions of agents.
2. Four months in a patient's life.
3. Stochastic simulations.

Simulations on GPUs.

1. FLAMEGPU and FLAMEGPU2 were used to generate optimised CUDA code.
2. 3000 time steps took up to 10 minutes.
3. Calibration took 40 days in total.



Hardware: 2 TITAN V GPUs, 1 TITAN XP GPU, and 1 TITAN RTX GPU.

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C

Clonal competition

Clonal composition.

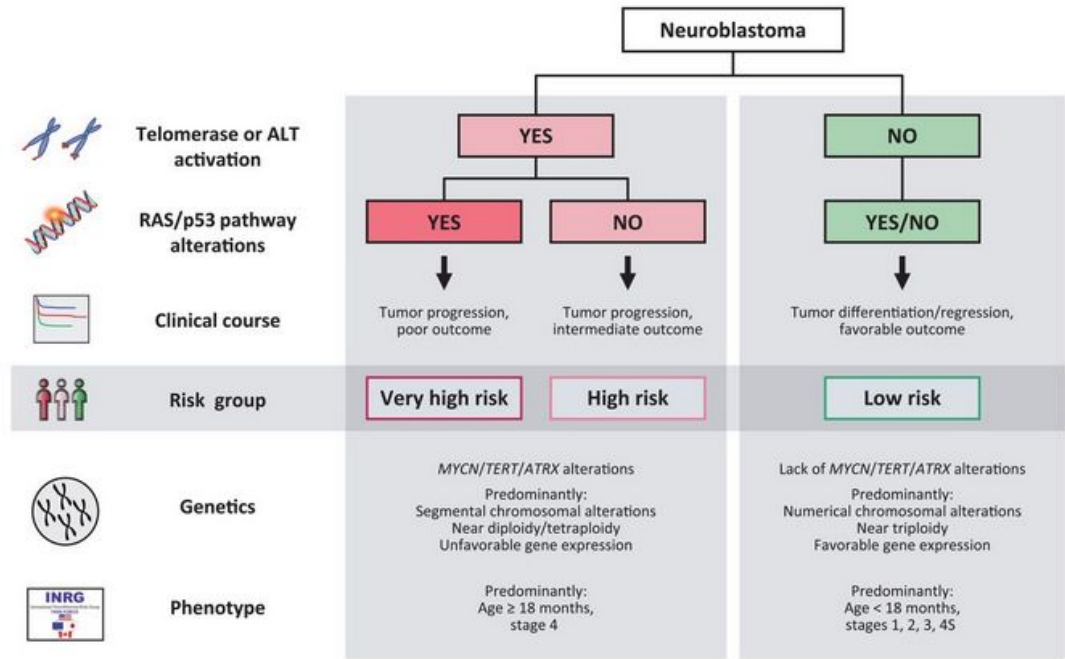
1. Four clones.
2. Each clone has six subclones.

Clones: MYCN amplification, TERT rearrangement, ATRX inactivation, and wild type.

Subclones: combinations of p53 inactivation and ALK activation.

Macroscopic features.

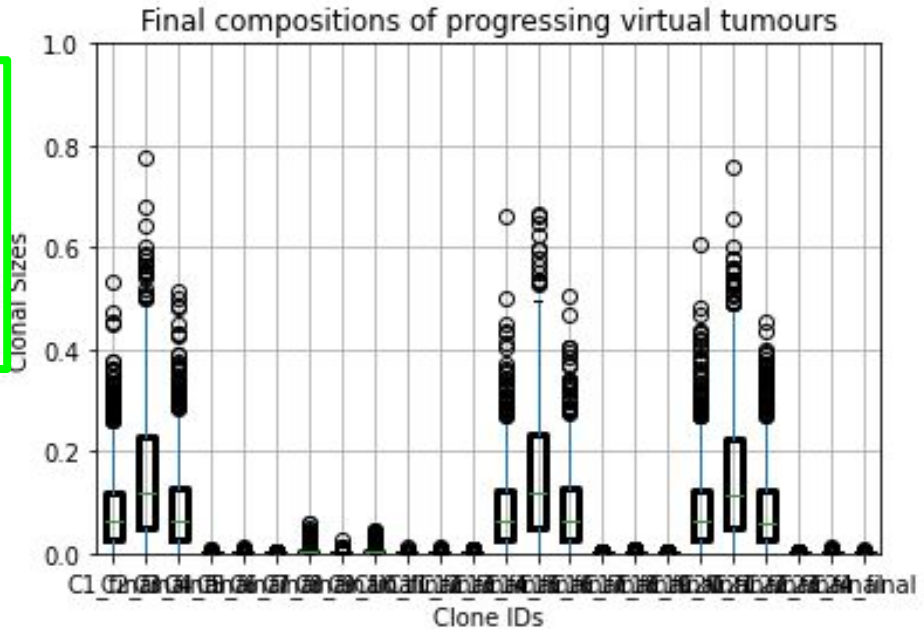
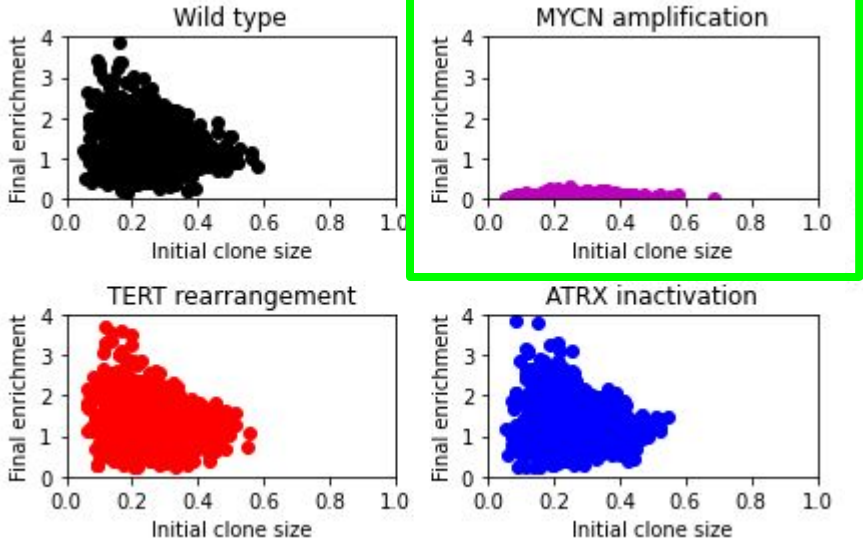
1. Oxygen level.
2. Abundance of Schwann cells.



Ackermann, Sandra, et al. "A mechanistic classification of clinical phenotypes in neuroblastoma." *Science* 362.6419 (2018): 1165-1170.

Created 1200 virtual tumours with arbitrary clonal compositions and macroscopic features.

Clonal competition



MYCN-amplified clone died!

The other three expanded similarly.

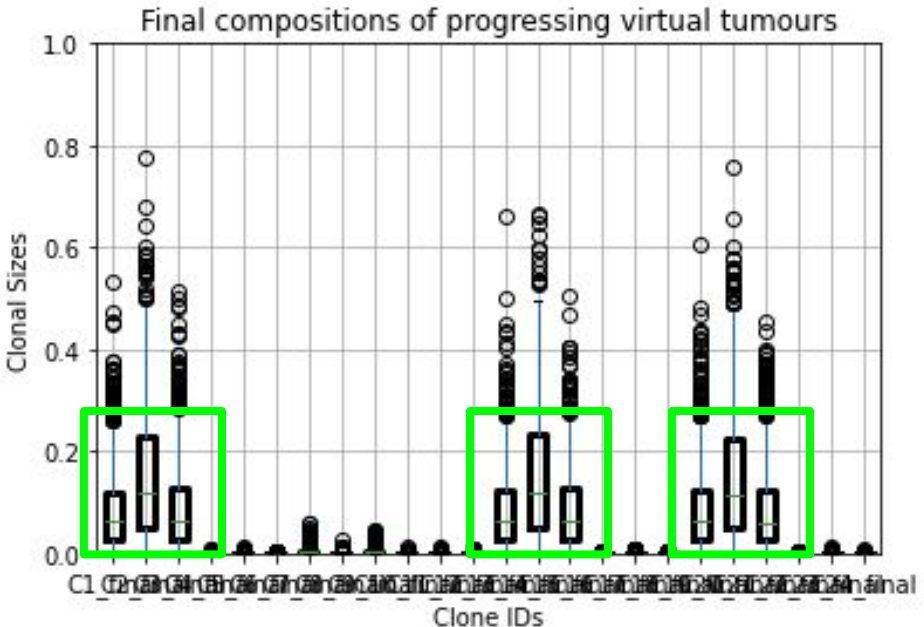
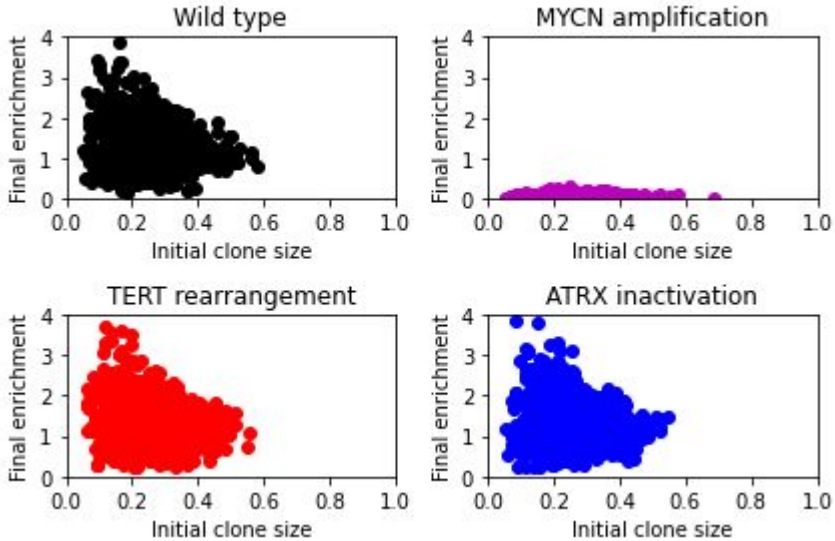
MA versus WT: p-value < 0.1 %

ANOVA: p-value > 25 %

1. Student's t-test.
2. Permutation test.

1. F-test.
2. Permutation test.

Clonal competition



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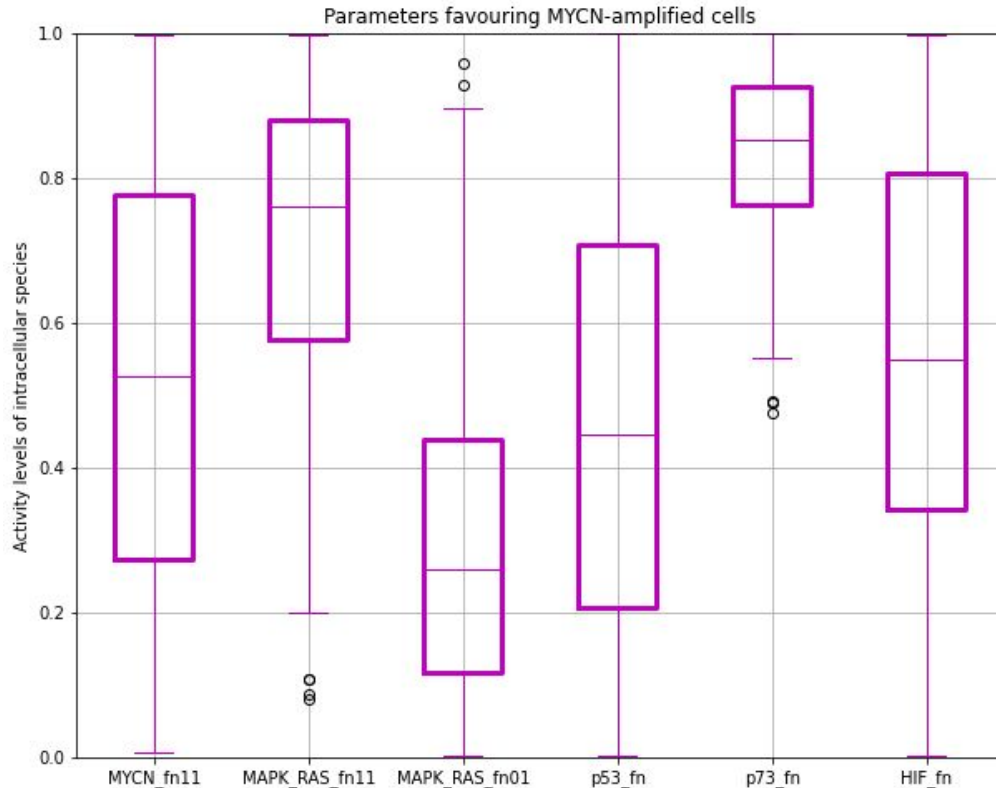
ANOVA: p-value > 25 %

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- 1. F-test.
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The nine growing subclones all had their p53 intact!

Clonal competition

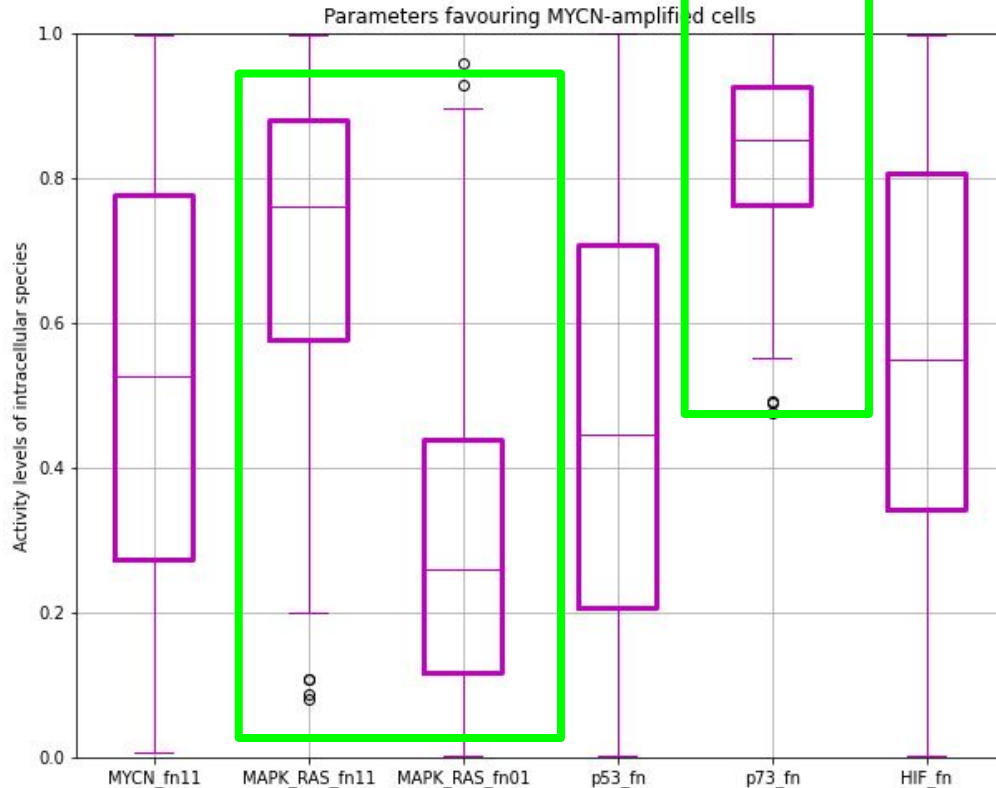


One virtual tumour. Fixed its clonal composition and macroscopic features.

Sensitivity analysis. 1000 combinations of gene expression levels, including MYCN, MAPK/RAS, p53, p73, and HIF.

283 cases where the MYCN-amplified clone expanded drastically.

Clonal competition



283 cases where the MYCN-amplified clone expanded drastically.

1. **p73**, which belongs to the same family, must be active enough to **compensate for the inactivated p53**.
2. **MYCN amplification** must boost MAPK/RAS signalling (**cell cycling**) more than **ALK activation**.

Objectives

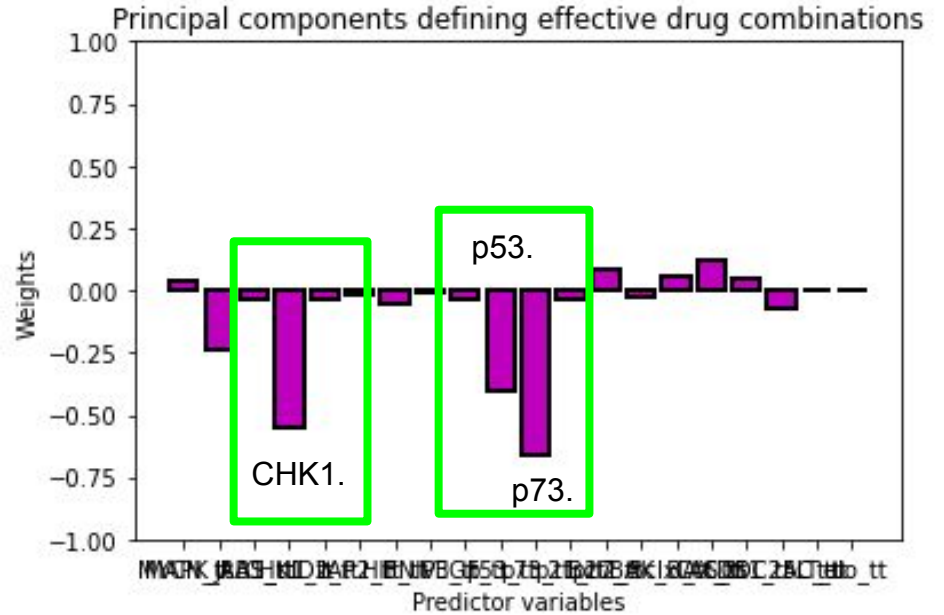
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Targeted therapies

Step 1. Created a virtual tumour with one large MYCN-amplified clone only.

Step 2. Chose gene expression levels favouring the MYCN-amplified clone.

Step 3. Tested 1000 combinations of drugs targeting the 20 gene products.



Weights of the first principal component.

Inhibiting CHK1, p53, and p73 shrinks the MYCN-amplified clone.

Consistent with sensitivity analysis, as CHK1 switches on p73.

20 gene products. Telomerase, ALT, MYCN, MAPK/RAS pathway, JAB1, CHK1, CDS1, CDC25C, ID2, IAP2, HIF, BNIP3, VEGF, p53, p73, p21, p27, Bcl-2/Bcl-xL, BAK/BAX, and CAS.

Conclusions

Built, calibrated, and validated the first multicellular model of neuroblastoma.

MYCN-Amplified clone requires p73 and enhanced cell cycling (MAPK/RAS signalling) to thrive.

Drugs targeting CHK1, p53, and p73 are effective against MYCN-amplified clone.